

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number	10 b10, 873	- .
Source:	1FW16	
Date Processed by STIC:	10-14-04	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE-LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04): 12 Colors of the U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04



IFW16

RAW SEQUENCE LISTING DATE: 10/14/2004 PATENT APPLICATION: US/10/010,873 TIME: 10:23:26

Input Set : A:\Sequence listing.txt
Output Set: N:\CRF4\10142004\J010873.raw

- 3 <110> APPLICANT: Babraham Bioscience Technologies Limited
- 5 <120> TITLE OF INVENTION: Recombinant Nematode Nicotinic Receptor and Uses
- 7 <130> FILE REFERENCE: 18396/2112
- C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/010,873
 - 10 <141> CURRENT FILING DATE: 2001-12-07
 - 12 <150> PRIOR APPLICATION NUMBER: PCT/GB00/02270
 - 13 <151> PRIOR FILING DATE: 2000-06-09
 - 15 <150> PRIOR APPLICATION NUMBER: GB9913248.2
 - 16 <151> PRIOR FILING DATE: 1999-06-09
 - 18 <160> NUMBER OF SEQ ID NOS: 4
 - 20 <170> SOFTWARE: PatentIn Ver. 3.1

Does Not Comply Corrected Diskette Needed

ERRORED SEQUENCES

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    126 <211> LENGTH: 513
    127 <212> TYPE: PRT
    128 <213> ORGANISM: Caenorhabditis elegans
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                  20
    137 Asp Leu Met Val Asn Tyr Asn Arg His Arg Arg Pro Ser Thr Ser Pro
                                40
              35
    140 Asn Lys Pro Leu Thr Ile Lys Leu Lys Leu Lys Leu Arg Leu Ser Gln
    143 Ile Ile Asp Val His Glu Ile Asp Gln Ile Met Thr Cys Ser Val Trp
                         70
                                           75
147
                      85
                                       90
  100
                                   105
                                                    110
    152 Pro Asp Ile Val Leu Tyr Asn Asn Ala Asp Ser Asn Tyr Asn Ile Thr
                               120
                                                 125
    155 Ile Ser Thr Lys Ala Thr Leu His Tyr Thr Gly Glu Val Thr Trp Glu
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    158 Pro Pro Ala Ile Phe Lys Ser Met Cys Gln Ile Asp Val Arg Trp Phe
                        150
    161 Pro Phe Asp Glu Gln Gln Cys His Leu Lys Phe Gly Ser Trp Thr Phe
                     165
                                      .170
    164 Ser Glu Asn Leu Leu Ser Val Glu Leu Asn Glu Pro Ser Leu Arg Tyr
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PATENT APPLICATION: US/10/010,873 TIME: 10:23:26

Input Set : A:\Sequence listing.txt
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	165				180					185					190		
E>	167	Glu	Glu	Glu	Ile	Asp	Glu	Lys	Gly	Ile	Ile	Asp	Asn	Val	Xaa	Xaa	Xaa
	168			195					200					205			
E>	170	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
	171		210					215					220				
E>	173	Xaa	Met	Ser	Arg	Val	Ala	Lys	Arg	Arg	Ala	Lys	Asn	Tyr	Pro	Ser	Cys
	174						230					235					240
	176	Cys	Pro	Gln	Ser	Ala	Tyr	Ile	Asp	Val		Tyr	Tyr	Leu	Gln		Arg
	177					245		_			250			_	-	255	
	179	Arg	Lys	Pro		Phe	Tyr	Thr	Val			Vai	Phe	Pro	Cys	Val	GIA
	180		_		260					265		•	n	~	270		41
		Ile	Ser		Leu	Thr	He	Leu		Pne	Tyr	Leu	Pro		Asp	ser	GIÅ
	183	_,		275	m1	•		~1 _	280	T 3.4	T	171	77.	285	TTID an	Tla	Dho
		GIU		vaı	Int	Leu	Cys	295	Ser	116	nea	var	300	Deu	Thr	116	FILE
	186	nha	290	T ON	Low	Thr	Glu		T.I.o.	Pro	λ1 =	Thr		Tle	Thr	T.611	Pro
		305	nea	ъęи	neu	1111	310	116	116	-10	nza	315	JUL			DCU	320
E>			Yaa	Yaa	Yaa	Kaa		Хаа	Xaa	Xaa	Хаа		Xaa	Xaa	Xaa	Xaa	
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E>			Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Leu	His	Phe	Arg	Thr	Pro	Thr
	195				340					345			•		350		
	197	Thr	His	Leu	Met	Pro	Asn	Trp	Val	Lys	Lys	Val	Phe	Leu	Lys	Trp	Leu
	198			355					360					365			_
	200	Pro	Lys	Leu	Leu	Phe	Met		Arg	Pro	Ile	Asp			Glu	Glu	Lys
	201	_	370		_	_	_	375	_	_		_	380		.	0	17.7
			Asp	Asp	rys	Lys		Pro	Lys	Asp	GIÅ		ire	Ala	Leu	Ser	va1 400
		385		***	N	17-7	390	N'am	17m3	Ø3.4	Nen	395	Tla		Asn	λla	
	205	HIS	ATA	UIR	Arg	405	Ser	ASII	vai	GLY	410	ABII	116	m. 9	Aon	415	****
B>		Tle	Agn	Asp	Thr		Gln	Lvs	Met	Tvr		Ser	Pro	Pro	Xaa		Xaa
	210				420		-	•		425		,	•		430		
E>		Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
	213			435				••	440					445			
E>	215	Xaa	Xaa	Xaa	Xaa	Ile	Asp	Glu	Asp	Trp	Lys	Tyr		Ala	Met	Val	Leu
	216		450					455					460				
		_	Arg	Leu	Phe	Leu		He	Phe	Ser	IIe		Cys	Phe	Val	GIÃ	
		465	~7.	¥1.	T	T	470	N 3 a	Dro	The	T 633	475	N cm	The	7-0	Gla	480 Pro
		vaı	TIE	116	гел	485	Arg	Ala	PIO	1111	490	ıyı	мер	1111	Arg	495	FIO
	222	۲l۵	han	Len	Gln		Δτα	Pro	Δla	Asn		Ser	Ala	Asn	Pro		Ser
	225	116	vob	Deu	500	-7-	**** 9		1124	505					510		
		Phe			•••												
			0 > S	EQ I	ON C	: 3											
				ENGT													
				YPE:													
	233	<21	3 > 0	RGAN:	ISM:	Cae	norha	abdi	tis (elega	ans		-				
				EQUE													
	236	Met	Met	Leu	Gly	Gly	Gly	Gly	Gly	Сув		Ala	Gly	Gly	Thr		Leu
•	237	1				5					10					15	

RAW SEQUENCE LISTING DATE: 10/14/2004 PATENT APPLICATION: US/10/010,873 TIME: 10:23:26

Input Set: A:\Sequence listing.txt
Output Set: N:\CRF4\10142004\J010873.raw

239 Gly Phe Leu Val Phe Leu Ala Val Ser Leu Arg Asn His Ser Thr Cys 242 Glu Asp Ile Asp Ala Glu Asp Arg Leu Met Val Asp Leu Phe Arg Gly 40 245 Tyr Asn Ser Leu Val Gln Pro Val Arg Asn Arg Ser Glu Leu Pro Met 248 Ile Val Lys Ile Gly Met Gln Leu Val Leu Leu Ile Asn Val Asp Glu 70 249 65 -> 251 Lys Glu Gln Val Met His Thr Asn Val Trp Leu Thr Met Lys Trp Xaa 85 90. 110 100 105 255 E--> 257 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Val Trp Leu Pro Asp Ile Val Leu Phe 120 258 115 260 Asn Asn Ala Asp Gly Asn Tyr Glu Val Ser Phe Met Cys Asn Val Leu 135 263 Ile Leu Ser Thr Gly Thr Val Leu Trp Val Pro Pro Ala Ile Tyr Lys 155 150 266 Ser Ser Cys Ile Ile Asp Val Glu Phe Phe Pro Phe Asp Asp Gln Leu 170 269 Cys Ser Leu Thr Phe Gly Ser Trp Thr Tyr Asn Arg Asp Glu Ile Lys 180 185 270 200 195 E--> 275 Xaa Xaa Xaa Xaa Xaa Xaa Met Asp Gly Pro Ala Val Leu Thr Ser Asp 215 278 Arg Ser Arg Ile Glu Phe Gln Ile Arg Ile Arg Lys Thr Leu Phe 279 225 230 235 281 Tyr Thr Val Val Leu Ile Leu Pro Thr Val Leu Met Ala Phe Leu Asn 245 284 Val Thr Val Phe Tyr Leu Pro Thr Ala Ser Gly Glu Lys Met Gly Leu 260 265 287 Thr Met Asn Val Leu Leu Ser Ile Val Val Phe Leu Leu Leu Val Ser 275 280 E--> 290 Lys Ile Leu Pro Pro Thr Ser Ser Ser Ile Pro Leu Xaa Xaa Xaa Xaa 310 315 E--> 296 Xaa Xaa Xaa Xaa Ile Tyr Phe Arg Ser Pro Ile Thr His Arg Leu Pro 325 330 299 Pro Trp Val Arg Lys Val Phe Leu Asp Ile Leu Pro Leu Leu Met Cys 340 345 302 Met Gln Arg Pro His Arg Lys Asn Val Ile Gln Arg Ser His Arg Arg 360 355 305 Leu Leu Glu Thr Gly Pro Ser Val Glu Glu Asn Pro Met Arg Ser Gly 375 380 370 308 Glu His His Pro Leu Cys Arg His Thr His Asn Gln Asp Ser Cys Arg 395 390 311 Arg Val Arg Ile Gln Ser Asp Glu Leu Asp Asp Glu Leu Ser Pro Glu

error

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Input Set: A:\Sequence listing.txt
Output Set: N:\CRF4\10142004\J010873.raw

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312
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315 420
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B--> 317 Xaa Xaa Xaa Xaa Xaa Xaa Phe Arg Asp Asp Trp Lys Phe Ile Ala
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                              440
    320 Ser Val Val Asp Arg Phe Leu Leu Tyr Gly Phe Phe Gly Ala Thr Val
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    333 <210> SEQ ID NO: 4
    334 <211> LENGTH: 493
    335 <212> TYPE: PRT
    336 <213> ORGANISM: Caenorhabditis elegans
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                                   25
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    345 Leu Met Val Asp Val Phe Arg Gly Tyr Asn Ser Leu Ile Gln Pro Val
    346 35
    348 Arg Asn Ser Ser Glu Leu Pro Leu Ile Val Lys Met Ala Leu Gln Leu
    351 Val Leu Leu Ile Asn Val Asp Glu Lys Asp Gln Val Met His Thr Asn
    352 65
                                                              journe error
E--> 354 Val Trp Leu Thr Leu Gln Trp Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                                            · 95
                                     90
105
                 100
    360 Trp Leu Pro Asp Ile Val Leu Phe Asn Asn Ala Asp Gly Asn Tyr Glu
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    361 115
                                             125
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    369 Phe Phe Pro Phe Asp Glu Gln Val Cys Thr Leu Val Phe Gly Ser Trp
                                     170
           . 165
 --> 372 Thr Tyr Asn Glu Asn Glu Ile Lys Leu Glu Phe Xaa Xaa Xaa Xaa
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    373
              180
200
        195
    378 Val Pro Ala Ser Leu Val Asn Lys Arg Ser Arg Ile Glu Phe Gln Val
                           215
    381 Arg Ile Arg Arg Lys Thr Leu Phe Tyr Thr Val Val Leu Ile Ile Pro
                       230
                                        235
    384 Thr Val Leu Met Ala Phe Leu Ser Met Ala Val Phe Phe Leu Pro Thr
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PATENT APPLICATION: US/10/010,873 TIME: 10:23:26

Input Set: A:\Sequence listing.txt
Output Set: N:\CRF4\10142004\J010873.raw

	385					245		•			250			٠		255		•
	387	Asp	Ser	Gly	Glu	Lys.	Ile	Thr	Leu		Ile	Ser	Val	Leu		Ser	Ile	
	388			•	260					265		_	_	_	270			
•		Val	Val		Leu	Leu	Leu	Val		Lys	He	Leu	Pro	285	Thr	ser	ser	
	391		-1-	275	W-4	¥	¥.	¥	280	¥	Vaa	Vaa	Vaa		Vaa	Yaa	Vaa	
E>	393	The	290	PIO	Add	Vaa	Add	295	Add.	naa	Add	Add	300	Add	Aud	1144	Auu	Same
E>		Xaa		Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Val	Tyr	Phe	Arg	Same
•	397						310					315					320	.
	399	Gly	Pro	Arg	Thr	His	Arg	Met	Pro	Gln	Trp	Val	Arg	۷al	Val		Leu	
	400					325					330		_	_	_	335	_	
		Gln	Phe	Leu		Lys	Leu	Val	Cys		Lys	Arg	Pro	Lys		Ala	ser	;
	403				340	**- *	1		61.	345		~1 m	Tous	Dro	350	T/= 1	G1v	
		GIU	Arg	355	ALA	vaı	Arg	ser	360	met	Ala	Gln	Leu	365	Gry	Vai	OLY	
	406	Gln	Dha		T.@11	Ser	Pro	Ser		His	His	Pro	Leu		Pro	Ser	Ala	
	409		370	****	u			375					380	-4 -				
				Arq	Thr	Thr	Thr	Ile	Arg	Asn	Thr	Ala	Ser	Asn	Glu	Thr	Ser	
	412	385	_				390					395					400	
E>	414	Ala	Tyr	Tyr	Pro	Leu	Ser	Thr	Asp	Xaa		Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	
	415					405	'				410					415	nh -	
E>			Xaa	Xaa			Xaa	Xaa	Xaa		Xaa	Xaa	хаа	Xaa	XAA	Aaa	Pne	
	418		3	3	420		Tire	Wal.	ב רג	425 Mat	Tla	Tla	λen	Δra		T.em	Leu	
	421		Asp	435		Бур	TYL	Val	440			110	p	445	200			•
	423	Tvr	Val			Glv	Ile	Thr			Gly	Thr	Cys	Gly	Ile	Leu	Phe	
	424		450					455		•	-		460	_				
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		_	Leu	Lys	Glu			Asp	Thr	Ala		Asn	Ile	Pro				
	430					485				-	490			•				

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/010,873

DATE: 10/14/2004 TIME: 10:23:27

Input Set : A:\Sequence listing.txt
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Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of <220> to <223> is MANDATORY if n's or Xaa's are present. in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

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Seg#:2; Xaa Pos. 214,215,216,217,218,219,220,221,222,223,224,225,322,323
Seg#:2; Xaa Pos. 324,325,326,327,328,329,330,331,332,333,334,335,336,337
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Seq#:4; Xaa Pos. 414,415,416,417,418,419,420,421,422,423,424,425,426,427
Seq#:4; Xaa Pos. 428,429,430,431
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VERIFICATION SUMMARY

DATE: 10/14/2004

PATENT APPLICATION: US/10/010,873

TIME: 10:23:27

Input Set : A:\Sequence listing.txt
Output Set: N:\CRF4\10142004\J010873.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number

L:146 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2

M:340 Repeated in SeqNo=2

L:251 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:3

M:340 Repeated in SeqNo=3

L:354 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:4

M:340 Repeated in SeqNo=4